



#10

-1-

SEQUENCE LISTING

<110> Chernajovsky, Yuti
Dreja, Hanna Stina
Adams, Gillian

<120> Latent Fusion Protein

<130> 0623.1000000

<140> US 09/756,283

<141> 2001-01-09

<160> 100

<170> PatentIn version 3.0

<210> 1

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<213> Artificial

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<211> 52

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23

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29

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<213> Artificial

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<223> Flexible linker

<400> 14

Gly Gly Gly Gly Ser
1 5

<210> 15

<211> 6

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<220>

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<400> 15

Pro Leu Gly Leu Trp Ala
1 5

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<223> Flexible portion

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<210> 17

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<400> 17

Pro Leu Gly Leu
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<220>

<223> Core of cleavage site

<400> 18

Pro Leu Gly Ile
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<210> 19

<211> 1376

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| atg | ccg | ccc | tcc | ggg | ctg | cgg | ctg | ctg | ccg | ctg | ctg | cta | ccg | ctg | ctg | |
| Met | Pro | Pro | Ser | Gly | Leu | Arg | Leu | Leu | Pro | Leu | Leu | Leu | Pro | Leu | Leu | |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | | |

48

| | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| tgg | cta | ctg | gtg | ctg | acg | cct | ggc | ccg | ccg | gcc | gcg | gga | cta | tcc | acc | |
| Trp | Leu | Leu | Val | Leu | Thr | Pro | Gly | Pro | Pro | Ala | Ala | Gly | Leu | Ser | Thr | |
| | | | 20 | | | | | 25 | | | | | 30 | | | |

96

| | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|--|
| tgc | aag | act | atc | gac | atg | gag | ctg | gtg | aag | cgg | aag | cgc | atc | gag | gcc | |
| Cys | Lys | Thr | Ile | Asp | Met | Glu | Leu | Val | Lys | Arg | Lys | Arg | Ile | Glu | Ala | |
| | | 35 | | | | | 40 | | | | | 45 | | | | |

144

| | | | | | | | | | | | | | | | | |
|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| atc Ile | cgc Arg 50 | ggc Gly | cag Gln | atc Ile | ctg Leu | tcc Ser 55 | aag Lys | ctg Leu | cgg Arg | ctc Leu | gcc Ala 60 | agc Ser | ccc Pro | ccg Pro | agc Ser | 192 |
| cag Gln 65 | ggg Gly | gag Glu | gtg Val | ccg Pro 70 | ccc Pro | ggc Gly | ccg Pro | ctg Leu | ccc Pro | gag Glu 75 | gcc Ala | gtg Val | ctc Leu | gcc Ala | ctg Leu 80 | 240 |
| tac Tyr | aac Asn | agc Ser | acc Thr | cgc Arg 85 | gac Asp | cgg Arg | gtg Val | gcc Ala | ggg Gly 90 | gag Glu | agt Ser | gca Ala | gaa Glu | ccg Pro 95 | gag Glu | 288 |
| ccc Pro | gag Glu | cct Pro | gag Glu 100 | gcc Ala | gac Asp | tac Tyr | tac Tyr | gcc Ala 105 | aag Lys | gag Glu | gtc Val | acc Thr | cgc Arg 110 | gtg Val | cta Leu | 336 |
| atg Met | gtg Val 115 | gaa Glu | acc Thr | cac His | aac Asn | gaa Glu | atc Ile 120 | tat Tyr | gac Asp | aag Lys | ttc Phe 125 | aag Lys | cag Gln | agt Ser | aca Thr | 384 |
| cac His 130 | agc Ser | ata Ile | tat Tyr | atg Met | ttc Phe | ttc Phe 135 | aac Asn | aca Thr | tca Ser | gag Glu | ctc Leu 140 | cga Arg | gaa Glu | gcg Ala | gta Val | 432 |
| cct Pro 145 | gaa Glu | ccc Pro | gtg Val | ttg Leu | ctc Leu 150 | tcc Ser | cgg Arg | gca Ala | gag Glu | ctg Leu 155 | cgt Arg | ctg Leu | ctg Leu | agg Arg | agg Arg 160 | 480 |
| ctc Leu | aag Lys | tta Leu | aaa Lys | gtg Val 165 | gag Glu | cag Gln | cac His | gtg Val | gag Glu 170 | ctg Leu | tac Tyr | cag Gln | aaa Lys | tac Tyr 175 | agc Ser | 528 |
| aac Asn | aat Asn | tcc Ser | tgg Trp 180 | cga Arg | tac Tyr | ctc Leu | agc Ser | aac Asn 185 | cgg Arg | ctg Leu | ctg Leu | gca Ala | ccc Pro 190 | agc Ser | gac Asp | 576 |
| tcg Ser | cca Pro | gag Glu 195 | tgg Trp | tta Leu | tct Ser | ttt Phe | gat Asp 200 | gtc Val | acc Thr | gga Gly | gtt Val | gtg Val 205 | cgg Arg | cag Gln | tgg Trp | 624 |
| ttg Leu 210 | agc Ser | cgt Arg | gga Gly | ggg Gly | gaa Glu | att Ile 215 | gag Glu | ggc Gly | ttt Phe | cgc Arg | ctt Leu 220 | agc Ser | gcc Ala | cac His | tgc Cys | 672 |
| tcc Ser 225 | tgt Cys | gac Asp | agc Ser | agg Arg | gat Asp 230 | aac Asn | aca Thr | ctg Leu | caa Gln | gtg Val 235 | gac Asp | atc Ile | aac Asn | ggg Gly | ttc Phe 240 | 720 |
| act Thr | acc Thr | ggc Gly | cgc Arg | cga Arg 245 | ggt Gly | gac Asp | ctg Leu | gcc Ala | acc Thr 250 | att Ile | cat His | ggc Gly | atg Met | aac Asn 255 | cgg Arg | 768 |
| cct Pro | ttc Phe | ctg Leu | ctt Leu 260 | ctc Leu | atg Met | gcc Ala | acc Thr 265 | ccg Pro | ctg Leu | gag Glu | agg Arg | gcc Ala | cag Gln 270 | cat His | ctg Leu | 816 |
| caa Gln | agc Ser | gaa Glu 275 | ttc Phe | ggg Gly | gga Gly | ggc Gly | gga Gly 280 | tcc Ser | ccg Pro | ctc Leu | ggg Gly | ctt Leu 285 | tgg Trp | gcg Ala | gga Gly | 864 |
| ggg Gly 290 | ggc Gly | tca Ser | gcg Ala | gcc Ala | gca Ala | atc Ile 295 | aac Asn | tat Tyr | aag Lys | cag Gln | ctc Leu 300 | cag Gln | ctc Leu | caa Gln | gaa Glu | 912 |
| agg Arg 305 | acg Thr | aac Asn | att Ile | cgg Arg | aaa Lys 310 | tgt Cys | cag Gln | gag Glu | ctc Leu 315 | ctg Leu | gag Glu | cag Gln | ctg Leu | aat Asn | gga Gly 320 | 960 |
| aag Lys | atc Ile | aac Asn | ctc Leu | acc Thr 325 | tac Tyr | agg Arg | gcg Ala | gac Asp | ttc Phe 330 | aag Lys | atc Ile | cct Pro | atg Met | gag Glu 335 | atg Met | 1008 |
| acg Thr | gag Glu | aag Lys | atg Met 340 | cag Gln | aag Lys | agt Ser | tac Tyr | act Thr 345 | gcc Ala | ttt Phe | gcc Ala | atc Ile | caa Gln 350 | gag Glu | atg Met | 1056 |
| ctc Leu | cag Gln | aat Asn | gtc Val | ttt Phe | ctt Leu | gtc Val | ttc Phe | aga Arg | aac Asn | aat Asn | ttc Phe | tcc Ser | agc Ser | act Thr | ggg Gly | 1104 |

| | 355 | | 360 | | 365 | |
|---|-----|-----|-----|-----|-----|------|
| tgg aat gag act att gtt gta cgt ctc ctg gat gaa ctc cac cag cag | | | | | | 1152 |
| Trp Asn Glu Thr Ile Val Val Arg Leu Leu Asp Glu Leu His Gln Gln | 370 | | 375 | | 380 | |
| aca gtg ttt ctg aag aca gta cta gag gaa aag caa gag gaa aga ttg | | | | | | 1200 |
| Thr Val Phe Leu Lys Thr Val Leu Glu Glu Lys Gln Glu Glu Arg Leu | 385 | | 390 | | 395 | 400 |
| acg tgg gag atg tcc tca act gct ctc cac ttg aag agc tat tac tgg | | | | | | 1248 |
| Thr Trp Glu Met Ser Ser Thr Ala Leu His Leu Lys Ser Tyr Tyr Trp | | 405 | | 410 | | 415 |
| agg gtg caa agg tac ctt aaa ctc atg aag tac aac agc tac gcc tgg | | | | | | 1296 |
| Arg Val Gln Arg Tyr Leu Lys Leu Met Lys Tyr Asn Ser Tyr Ala Trp | | 420 | | 425 | | 430 |
| atg gtg gtc cga gca gag atc ttc agg aac ttt ctc atc att cga aga | | | | | | 1344 |
| Met Val Val Arg Ala Glu Ile Phe Arg Asn Phe Leu Ile Ile Arg Arg | | 435 | | 440 | | 445 |
| ctt acc aga aac ttc caa aac tga tctagacc | | | | | | 1376 |
| Leu Thr Arg Asn Phe Gln Asn | 450 | | 455 | | | |

<210> 20

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<213> Artificial

<220>

<223> LAP-mIFN β construct

<400> 20

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|---|-----|-----|-----|----|
| Met Pro Pro Ser Gly Leu Arg Leu Leu Pro Leu Leu Leu Pro Leu Leu | 1 | 5 | 10 | 15 |
| Trp Leu Leu Val Leu Thr Pro Gly Pro Pro Ala Ala Gly Leu Ser Thr | 20 | 25 | 30 | |
| Cys Lys Thr Ile Asp Met Glu Leu Val Lys Arg Lys Arg Ile Glu Ala | 35 | 40 | 45 | |
| Ile Arg Gly Gln Ile Leu Ser Lys Leu Arg Leu Ala Ser Pro Pro Ser | 50 | 55 | 60 | |
| Gln Gly Glu Val Pro Pro Gly Pro Leu Pro Glu Ala Val Leu Ala Leu | 65 | 70 | 75 | 80 |
| Tyr Asn Ser Thr Arg Asp Arg Val Ala Gly Glu Ser Ala Glu Pro Glu | 85 | 90 | 95 | |
| Pro Glu Pro Glu Ala Asp Tyr Tyr Ala Lys Glu Val Thr Arg Val Leu | 100 | 105 | 110 | |
| Met Val Glu Thr His Asn Glu Ile Tyr Asp Lys Phe Lys Gln Ser Thr | 115 | 120 | 125 | |
| His Ser Ile Tyr Met Phe Phe Asn Thr Ser Glu Leu Arg Glu Ala Val | 130 | 135 | 140 | |

Pro Glu Pro Val Leu Leu Ser Arg Ala Glu Leu Arg Leu Leu Arg Arg
 145 150 155 160
 Leu Lys Leu Lys Val Glu Gln His Val Glu Leu Tyr Gln Lys Tyr Ser
 165 170 175
 Asn Asn Ser Trp Arg Tyr Leu Ser Asn Arg Leu Leu Ala Pro Ser Asp
 180 185 190
 Ser Pro Glu Trp Leu Ser Phe Asp Val Thr Gly Val Val Arg Gln Trp
 195 200 205
 Leu Ser Arg Gly Gly Glu Ile Glu Gly Phe Arg Leu Ser Ala His Cys
 210 215 220
 Ser Cys Asp Ser Arg Asp Asn Thr Leu Gln Val Asp Ile Asn Gly Phe
 225 230 235 240
 Thr Thr Gly Arg Arg Gly Asp Leu Ala Thr Ile His Gly Met Asn Arg
 245 250 255
 Pro Phe Leu Leu Leu Met Ala Thr Pro Leu Glu Arg Ala Gln His Leu
 260 265 270
 Gln Ser Glu Phe Gly Gly Gly Gly Ser Pro Leu Gly Leu Trp Ala Gly
 275 280 285
 Gly Gly Ser Ala Ala Ala Ile Asn Tyr Lys Gln Leu Gln Leu Gln Glu
 290 295 300
 Arg Thr Asn Ile Arg Lys Cys Gln Glu Leu Leu Glu Gln Leu Asn Gly
 305 310 315 320
 Lys Ile Asn Leu Thr Tyr Arg Ala Asp Phe Lys Ile Pro Met Glu Met
 325 330 335
 Thr Glu Lys Met Gln Lys Ser Tyr Thr Ala Phe Ala Ile Gln Glu Met
 340 345 350
 Leu Gln Asn Val Phe Leu Val Phe Arg Asn Asn Phe Ser Ser Thr Gly
 355 360 365
 Trp Asn Glu Thr Ile Val Val Arg Leu Leu Asp Glu Leu His Gln Gln
 370 375 380
 Thr Val Phe Leu Lys Thr Val Leu Glu Glu Lys Gln Glu Glu Arg Leu
 385 390 395 400
 Thr Trp Glu Met Ser Ser Thr Ala Leu His Leu Lys Ser Tyr Tyr Trp
 405 410 415
 Arg Val Gln Arg Tyr Leu Lys Leu Met Lys Tyr Asn Ser Tyr Ala Trp
 420 425 430
 Met Val Val Arg Ala Glu Ile Phe Arg Asn Phe Leu Ile Ile Arg Arg
 435 440 445
 Leu Thr Arg Asn Phe Gln Asn


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450
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<211> 1352
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<220>
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<220>
<221> CDS
<222> (1)..(1344)

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Met Asn Asn Arg Trp Ile Leu His Ala Ala Phe Leu Leu Cys Phe Ser
1 5 10 15
acc aca gcc ctc tcc atc aac tat aag cag ctc cag ctc caa gaa agg 96
Thr Thr Ala Leu Ser Ile Asn Tyr Lys Gln Leu Gln Leu Gln Glu Arg
20 25 30
acg aac att cgg aaa tgt cag gag ctc ctg gag cag ctg aat gga aag 144
Thr Asn Ile Arg Lys Cys Gln Glu Leu Leu Glu Gln Leu Asn Gly Lys
35 40 45
atc aac ctc acc tac agg gcg gac ttc aag atc cct atg gag atg acg 192
Ile Asn Leu Thr Tyr Arg Ala Asp Phe Lys Ile Pro Met Glu Met Thr
50 55 60
gag aag atg cag aag agt tac act gcc ttt gcc atc caa gag atg ctc 240
Glu Lys Met Gln Lys Ser Tyr Thr Ala Phe Ala Ile Gln Glu Met Leu
65 70 75 80
cag aat gtc ttt ctt gtc ttc aga aac aat ttc tcc agc act ggg tgg 288
Gln Asn Val Phe Leu Val Phe Arg Asn Asn Phe Ser Ser Thr Gly Trp
85 90 95
aat gag act att gtt gta cgt ctc ctg gat gaa ctc cac cag cag aca 336
Asn Glu Thr Ile Val Val Arg Leu Leu Asp Glu Leu His Gln Gln Thr
100 105 110
gtg ttt ctg aag aca gta cta gag gaa aag caa gag gaa aga ttg acg 384
Val Phe Leu Lys Thr Val Leu Glu Glu Lys Gln Glu Glu Arg Leu Thr
115 120 125
tgg gag atg tcc tca act gct ctc cac ttg aag agc tat tac tgg agg 432
Trp Glu Met Ser Ser Thr Ala Leu His Leu Lys Ser Tyr Tyr Trp Arg
130 135 140
gtg caa agg tac ctt aaa ctc atg aag tac aac agc tac gcc tgg atg 480
Val Gln Arg Tyr Leu Lys Leu Met Lys Tyr Asn Ser Tyr Ala Trp Met
145 150 155 160
gtg gtc cga gca gag atc ttc agg aac ttt ctc atc att cga aga ctt 528
Val Val Arg Ala Glu Ile Phe Arg Asn Phe Leu Ile Ile Arg Arg Leu
165 170 175
acc aga aac ttc caa aac gaa ttc ggg gga ggc gga tcc ccg ctc ggg 576
Thr Arg Asn Phe Gln Asn Glu Phe Gly Gly Gly Gly Ser Pro Leu Gly
180 185 190
ctt tgg gcg gga ggg ggc tca gcg gcc gca cta tcc acc tgc aag act 624
Leu Trp Ala Gly Gly Gly Ser Ala Ala Leu Ser Thr Cys Lys Thr
195 200 205

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| | | | | | | | | | | | | | | | | |
|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|------|
| atc Ile | gac Asp 210 | atg Met | gag Glu | ctg Leu | gtg Val | aag Lys 215 | cgg Arg | aag Lys | cgc Arg | atc Ile | gag Glu 220 | gcc Ala | atc Ile | cgc Arg | ggc Gly | 672 |
| cag Gln 225 | atc Ile | ctg Leu | tcc Ser | aag Lys | ctg Leu 230 | cgg Arg | ctc Leu | gcc Ala | agc Ser | ccc Pro 235 | ccg Pro | agc Ser | cag Gln | ggg Gly | gag Glu 240 | 720 |
| gtg Val | ccg Pro | ccc Pro | ggc Gly | ccg Pro 245 | ctg Leu | ccc Pro | gag Glu | gcc Ala | gtg Val 250 | ctc Leu | gcc Ala | ctg Leu | tac Tyr | aac Asn 255 | agc Ser | 768 |
| acc Thr | cgc Arg | gac Asp | cgg Arg 260 | gtg Val | gcc Ala | ggg Gly | gag Glu | agt Ser 265 | gca Ala | gaa Glu | ccg Pro | gag Glu | ccc Pro 270 | gag Glu | cct Pro | 816 |
| gag Glu | gcc Ala | gac Asp 275 | tac Tyr | tac Tyr | gcc Ala | aag Lys | gag Glu 280 | gtc Val | acc Thr | cgc Arg | gtg Val | cta Leu 285 | atg Met | gtg Val | gaa Glu | 864 |
| acc Thr | cac His 290 | aac Asn | gaa Glu | atc Ile | tat Tyr | gac Asp 295 | aag Lys | ttc Phe | aag Lys | cag Gln | agt Ser 300 | aca Thr | cac His | agc Ser | ata Ile | 912 |
| tat Tyr 305 | atg Met | ttc Phe | ttc Phe | aac Asn | aca Thr 310 | tca Ser | gag Glu | ctc Leu | cga Arg | gaa Glu 315 | gcg Ala | gta Val | cct Pro | gaa Glu | ccc Pro 320 | 960 |
| gtg Val | ttg Leu | ctc Leu | tcc Ser | cgg Arg 325 | gca Ala | gag Glu | ctg Leu | cgt Arg | ctg Leu 330 | ctg Leu | agg Arg | agg Arg | ctc Leu | aag Lys 335 | tta Leu | 1008 |
| aaa Lys | gtg Val | gag Glu | cag Gln 340 | cac His | gtg Val | gag Glu | ctg Leu | tac Tyr 345 | cag Gln | aaa Lys | tac Tyr | agc Ser | aac Asn 350 | aat Asn | tcc Ser | 1056 |
| tgg Trp | cga Arg | tac Tyr 355 | ctc Leu | agc Ser | aac Asn | cgg Arg | ctg Leu 360 | ctg Leu | gca Ala | ccc Pro | agc Ser | gac Asp 365 | tcg Ser | cca Pro | gag Glu | 1104 |
| tgg Trp 370 | tta Leu | tct Ser | ttt Phe | gat Asp | gtc Val | acc Thr 375 | gga Gly | gtt Val | gtg Val | cgg Arg | cag Gln 380 | tgg Trp | ttg Leu | agc Ser | cgt Arg | 1152 |
| gga Gly 385 | ggg Gly | gaa Glu | att Ile | gag Glu | ggc Gly 390 | ttt Phe | cgc Arg | ctt Leu | agc Ser | gcc Ala 395 | cac His | tgc Cys | tcc Ser | tgt Cys | gac Asp 400 | 1200 |
| agc Ser | agg Arg | gat Asp | aac Asn | aca Thr 405 | ctg Leu | caa Gln | gtg Val | gac Asp | atc Ile 410 | aac Asn | ggg Gly | ttc Phe | act Thr | acc Thr 415 | ggc Gly | 1248 |
| cgc Arg | cga Arg | ggt Gly | gac Asp 420 | ctg Leu | gcc Ala | acc Thr | att Ile | cat His 425 | ggc Gly | atg Met | aac Asn | cgg Arg | cct Pro 430 | ttc Phe | ctg Leu | 1296 |
| ctt Leu | ctc Leu | atg Met 435 | gcc Ala | acc Thr | cgg Pro | ctg Leu | gag Glu 440 | agg Arg | gcc Ala | cag Gln | cat His | ctg Leu 445 | caa Gln | agc Ser | tga | 1344 |
| tctagacc | | | | | | | | | | | | | | | | 1352 |

<210> 22

<211> 447

<212> PRT

<213> Artificial

<220>

<223> mIFN β -LAP construct

<400> 22

Met Asn Asn Arg Trp Ile Leu His Ala Ala Phe Leu Leu Cys Phe Ser
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20 25 30

Thr Asn Ile Arg Lys Cys Gln Glu Leu Leu Glu Gln Leu Asn Gly Lys
35 40 45

Ile Asn Leu Thr Tyr Arg Ala Asp Phe Lys Ile Pro Met Glu Met Thr
50 55 60

Glu Lys Met Gln Lys Ser Tyr Thr Ala Phe Ala Ile Gln Glu Met Leu
65 70 75 80

Gln Asn Val Phe Leu Val Phe Arg Asn Asn Phe Ser Ser Thr Gly Trp
85 90 95

Asn Glu Thr Ile Val Val Arg Leu Leu Asp Glu Leu His Gln Gln Thr
100 105 110

Val Phe Leu Lys Thr Val Leu Glu Glu Lys Gln Glu Glu Arg Leu Thr
115 120 125

Trp Glu Met Ser Ser Thr Ala Leu His Leu Lys Ser Tyr Tyr Trp Arg
130 135 140

Val Gln Arg Tyr Leu Lys Leu Met Lys Tyr Asn Ser Tyr Ala Trp Met
145 150 155 160

Val Val Arg Ala Glu Ile Phe Arg Asn Phe Leu Ile Ile Arg Arg Leu
165 170 175

Thr Arg Asn Phe Gln Asn Glu Phe Gly Gly Gly Gly Ser Pro Leu Gly
180 185 190

Leu Trp Ala Gly Gly Gly Ser Ala Ala Ala Leu Ser Thr Cys Lys Thr
195 200 205

Ile Asp Met Glu Leu Val Lys Arg Lys Arg Ile Glu Ala Ile Arg Gly
210 215 220

Gln Ile Leu Ser Lys Leu Arg Leu Ala Ser Pro Pro Ser Gln Gly Glu
225 230 235 240

Val Pro Pro Gly Pro Leu Pro Glu Ala Val Leu Ala Leu Tyr Asn Ser
245 250 255

Thr Arg Asp Arg Val Ala Gly Glu Ser Ala Glu Pro Glu Pro Glu Pro
260 265 270

Glu Ala Asp Tyr Tyr Ala Lys Glu Val Thr Arg Val Leu Met Val Glu
275 280 285

Thr His Asn Glu Ile Tyr Asp Lys Phe Lys Gln Ser Thr His Ser Ile
290 295 300

Tyr Met Phe Phe Asn Thr Ser Glu Leu Arg Glu Ala Val Pro Glu Pro
305 310 315 320

Val Leu Leu Ser Arg 325 Ala Glu Leu Arg Leu 330 Leu Arg Arg Leu Lys 335 Leu
 Lys Val Glu Gln 340 His Val Glu Leu Tyr 345 Gln Lys Tyr Ser Asn 350 Asn Ser
 Trp Arg Tyr 355 Leu Ser Asn Arg Leu 360 Leu Ala Pro Ser Asp 365 Ser Pro Glu
 Trp Leu 370 Ser Phe Asp Val Thr 375 Gly Val Val Arg Gln 380 Trp Leu Ser Arg
 Gly 385 Gly Glu Ile Glu Gly 390 Phe Arg Leu Ser Ala 395 His Cys Ser Cys Asp 400
 Ser Arg Asp Asn Thr 405 Leu Gln Val Asp Ile 410 Asn Gly Phe Thr Thr 415 Gly
 Arg Arg Gly Asp 420 Leu Ala Thr Ile His 425 Gly Met Asn Arg Pro 430 Phe Leu
 Leu Leu Met 435 Ala Thr Pro Leu Glu 440 Arg Ala Gln His Leu 445 Gln Ser

<210> 23

<211> 390

<212> PRT

<213> Homo sapiens

<400> 23

Met Pro Pro Ser Gly 5 Leu Arg Leu Leu Pro 10 Leu Leu Leu Pro Leu 15 Leu
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 Cys Lys Thr 35 Ile Asp Met Glu 40 Leu Val Lys Arg Lys 45 Arg Ile Glu Ala
 Ile Arg 50 Gly Gln Ile Leu Ser 55 Lys Leu Arg Leu Ala 60 Ser Pro Pro Ser
 Gln 65 Gly Glu Val Pro 70 Pro Gly Pro Leu Pro Glu 75 Ala Val Leu Ala Leu 80
 Tyr Asn Ser Thr Arg 85 Asp Arg Val Ala Gly 90 Glu Ser Ala Glu 95 Pro Glu
 Pro Glu Pro Glu 100 Ala Asp Tyr Tyr Ala 105 Lys Glu Val Thr Arg 110 Val Leu
 Met Val Glu 115 Thr His His Glu Ile 120 Tyr Asp Lys Phe Lys 125 Gln Ser Thr
 His Ser 130 Thr Tyr Met Phe Phe 135 Asn Ile Ser Glu Leu 140 Arg Glu Ala Val
 Pro 145 Glu Pro Val Leu Leu 150 Ser Arg Ala Glu Leu 155 Arg Leu Leu Arg Leu 160
 Lys Leu Lys Val Glu 165 Gln His Val Glu Leu 170 Tyr Gln Lys Tyr Ser 175 Asn

Asn Ser Trp Arg Tyr Leu Ser Asn Arg Leu Leu Ala Pro Ser Asp Ser
180 185 190
Pro Glu Trp Leu Ser Phe Asp Val Thr Gly Val Val Arg Gln Trp Leu
195 200 205
Ser Arg Gly Gly Glu Ile Glu Gly Phe Arg Leu Ser Ala His Cys Ser
210 215 220
Cys Asp Ser Arg Asp Asn Thr Leu Gln Val Asp Ile Asn Gly Phe Thr
225 230 235 240
Thr Gly Arg Arg Gly Asp Leu Ala Thr Ile His Gly Met Asn Arg Pro
245 250 255
Phe Leu Leu Leu Met Ala Thr Pro Leu Glu Arg Ala Gln His Leu Gln
260 265 270
Ser Ser Arg His Arg Arg Ala Leu Asp Thr Asn Tyr Cys Phe Ser Ser
275 280 285
Thr Glu Lys Asn Cys Cys Val Arg Gln Leu Tyr Ile Asp Phe Arg Lys
290 300
Asp Leu Gly Trp Lys Trp Ile His Glu Pro Lys Gly Tyr His Ala Asn
305 310 315 320
Phe Cys Leu Gly Pro Cys Pro Tyr Ile Trp Ser Leu Asp Thr Gln Tyr
325 330 335
Ser Lys Val Leu Ala Leu Tyr Asn Gln His Asn Pro Gly Ala Ser Ala
340 345 350
Ala Pro Cys Cys Val Pro Gln Ala Leu Glu Pro Leu Pro Ile Val Tyr
355 360 365
Tyr Val Gly Arg Lys Pro Lys Val Glu Gln Leu Ser Asn Met Ile Val
370 375 380
Arg Ser Cys Lys Cys Ser
385 390

<210> 24

<211> 414

<212> PRT

<213> Homo sapiens

<400> 24

Met His Tyr Cys Val Leu Ser Ala Phe Leu Ile Leu His Leu Val Thr
1 5 10 15
Val Ala Leu Ser Leu Ser Thr Cys Ser Thr Leu Asp Met Gln Gln Phe
20 25 30
Met Arg Lys Arg Ile Glu Ala Ile Arg Gly Gln Ile Leu Ser Lys Leu
35 40 45
Lys Leu Thr Ser Pro Pro Glu Asp Tyr Pro Glu Pro Glu Glu Val Pro
50 55 60
Pro Glu Val Ile Ser Ile Tyr Asn Ser Thr Arg Asp Leu Leu Gln Glu
65 70 75 80
Lys Ala Ser Arg Arg Ala Ala Ala Cys Glu Arg Glu Arg Ser Asp Glu
85 90 95
Glu Tyr Tyr Ala Lys Glu Val Tyr Lys Ile Asp Met Pro Pro Phe Phe
100 105 110
Pro Ser Glu Asn Ala Ile Pro Pro Thr Phe Tyr Arg Pro Tyr Phe Arg
115 120 125

Ile Val Arg Phe Asp Val Ser Ala Met Glu Lys Asn Ala Ser Asn Leu
130 135 140
Val Lys Ala Glu Phe Arg Val Phe Arg Leu Gln Asn Pro Lys Ala Arg
145 150 155 160
Val Pro Glu Gln Arg Ile Glu Leu Tyr Gln Ile Leu Lys Ser Lys Asp
165 170 175
Leu Ile Ser Pro Thr Gln Arg Tyr Ile Asp Ser Lys Val Val Lys Thr
180 185 190
Arg Ala Glu Gly Glu Trp Leu Ser Phe Asp Val Thr Asp Ala Val His
195 200 205
Glu Trp Leu His His Lys Asp Arg Asn Leu Gly Phe Lys Ile Ser Leu
210 215 220
His Cys Pro Cys Cys Thr Phe Val Pro Ser Asn Asn Tyr Ile Ile Pro
225 230 235 240
Asn Lys Ser Glu Glu Leu Glu Ala Arg Phe Ala Gly Ile Asp Gly Ile
245 250 255
Ser Thr Tyr Thr Ser Gly Asp Gln Lys Thr Ile Lys Ser Thr Arg Lys
260 265 270
Lys Asn Ser Gly Lys Thr Pro His Leu Leu Leu Met Leu Leu Pro Ser
275 280 285
Tyr Arg Leu Glu Ser Gln Gln Thr Asn Arg Arg Lys Lys Arg Ala Leu
290 295 300
Asp Ala Ala Tyr Cys Phe Arg Asn Val Gln Asp Asn Cys Cys Leu Arg
305 310 315 320
Pro Leu Tyr Ile Asp Phe Lys Arg Asp Leu Gly Trp Lys Trp Ile His
325 330 335
Glu Pro Lys Gly Tyr Asn Ala Asn Phe Cys Ala Gly Ala Cys Pro Tyr
340 345 350
Leu Trp Ser Ser Asp Thr Gln His Ser Arg Val Leu Ser Leu Tyr Asn
355 360 365
Thr Glu Asn Pro Glu Ala Ser Ala Ser Pro Cys Cys Val Ser Gln Asp
370 375 380
Leu Glu Pro Leu Thr Ile Leu Tyr Tyr Ile Gly Lys Ile Pro Lys Ile
385 390 395 400
Glu Gln Leu Ser Asn Met Ile Val Lys Ser Cys Lys Cys Ser
405 410

<210> 25

<211> 412

<212> PRT

<213> Homo sapiens

<400> 25

Met Lys Met His Leu Gln Arg Ala Leu Val Val Leu Ala Leu Leu His
1 5 10 15
Phe Ala Thr Val Ser Leu Ser Leu Ser Thr Cys Thr Thr Leu Asp Phe
20 25 30
Gly His Ile Lys Lys Lys Arg Val Glu Ala Ile Arg Gly Gln Ile Leu
35 40 45
Ser Lys Leu Arg Leu Thr Ser Pro Pro Glu Pro Thr Val Met Thr His
50 55 60

Val Pro Tyr Gln Val Leu Ala Leu Tyr Asn Ser Thr Arg Glu Leu Leu
65 70 75 80
Glu Glu His Gly Glu Arg Lys Glu Glu Gly Cys Thr Gln Glu Asn Thr
85 90 95
Glu Ser Glu Tyr Tyr Ala Lys Glu Ile His Lys Phe Asp Met Ile Gln
100 105 110
Gly Leu Ala Glu His Asn Glu Leu Ala Val Cys Pro Lys Gly Ile Thr
115 120 125
Ser Lys Val Phe Arg Phe Asn Val Ser Ser Val Glu Lys Asn Arg Thr
130 135 140
Asn Leu Phe Arg Ala Glu Phe Arg Val Leu Arg Val Pro Asn Pro Ser
145 150 155 160
Ser Lys Arg Asn Glu Gln Arg Ile Glu Leu Phe Gln Ile Leu Arg Pro
165 170 175
Asp Glu His Ile Ala Lys Gln Arg Tyr Ile Gly Gly Lys Asn Leu Pro
180 185 190
Thr Arg Gly Thr Ala Glu Trp Leu Ser Phe Asp Val Thr Asp Thr Val
195 200 205
Arg Glu Trp Leu Leu Arg Arg Glu Ser Asn Leu Gly Leu Glu Ile Ser
210 215 220
Ile His Cys Pro Cys His Thr Phe Gln Pro Asn Gly Asp Ile Leu Glu
225 230 235 240
Asn Ile His Glu Val Met Glu Ile Lys Phe Lys Gly Val Asp Asn Glu
245 250 255
Asp Asp His Gly Arg Gly Asp Leu Gly Arg Leu Lys Lys Gln Lys Asp
260 265 270
Asn Asn Asn Pro His Leu Ile Leu Met Met Ile Pro Pro His Arg Leu
275 280 285
Asp Asn Pro Gly Gln Gly Gly Gln Arg Lys Lys Arg Ala Leu Asp Ile
290 295 300
Asn Tyr Cys Phe Arg Asn Leu Glu Glu Asn Cys Cys Val Arg Pro Leu
305 310 315 320
Tyr Ile Asp Phe Arg Gln Asp Leu Gly Trp Lys Trp Val His Glu Pro
325 330 335
Lys Gly Tyr Tyr Ala Asn Phe Cys Ser Gly Pro Cys Pro Tyr Leu Arg
340 345 350
Ser Ala Asp Thr Thr His Ser Thr Val Leu Gly Leu Tyr Asn Thr Leu
355 360 365
Asn Pro Glu Ala Ser Ala Ser Pro Cys Cys Val Pro Gln Asp Leu Glu
370 375 380
Pro Leu Thr Ile Leu Tyr Tyr Val Gly Arg Thr Pro Lys Val Glu Gln
385 390 395 400
Leu Ser Asn Met Val Val Lys Ser Cys Lys Cys Ser
405 410

<210> 26

<211> 304

<212> PRT

<213> Gallus domesticus

<400> 26

Met Asp Pro Met Ser Ile Gly Pro Lys Ser Cys Gly Gly Ser Pro Trp
1 5 10 15
Arg Pro Pro Gly Thr Ala Pro Trp Ser Ile Gly Ser Arg Arg Ala Thr
20 25 30
Ala Ser Ser Ser Cys Ser Thr Ser Arg Val Arg Ala Glu Val Gly
35 40 45
Gly Arg Ala Leu Leu His Arg Ala Glu Leu Arg Met Leu Arg Gln Lys
50 55 60
Ala Ala Ala Asp Ser Ala Gly Thr Glu Gln Arg Leu Glu Leu Tyr Gln
65 70 75
Gly Tyr Gly Asn Ala Ser Trp Arg Tyr Leu His Gly Arg Ser Val Arg
85 90 95
Ala Thr Ala Asp Asp Glu Trp Leu Ser Phe Asp Val Thr Asp Ala Val
100 105 110
His Gln Trp Leu Ser Gly Ser Glu Leu Leu Gly Val Phe Lys Leu Ser
115 120 125
Val His Cys Pro Cys Glu Met Gly Pro Gly His Ala Asp Glu Met Arg
130 135 140
Ile Ser Ile Glu Gly Phe Glu Gln Gln Arg Gly Asp Met Gln Ser Ile
145 150 155
Ala Lys Lys His Arg Arg Val Pro Tyr Val Leu Ala Met Ala Leu Pro
165 170 175
Ala Glu Arg Ala Asn Glu Leu His Ser Ala Arg Arg Arg Arg Asp Leu
180 185 190
Asp Thr Asp Tyr Cys Phe Gly Pro Gly Thr Asp Glu Lys Asn Cys Cys
195 200 205
Val Arg Pro Leu Tyr Ile Asp Phe Arg Lys Asp Leu Gln Trp Lys Trp
210 215 220
Ile His Glu Pro Lys Gly Tyr Met Ala Asn Phe Cys Met Gly Pro Cys
225 230 235
Pro Tyr Ile Trp Ser Ala Asp Thr Gln Tyr Ile Lys Val Leu Ala Leu
245 250 255
Tyr Asn Gln Asn Asn Pro Gly Ala Ser Ala Ala Pro Cys Cys Val Pro
260 265 270
Gln Ile Leu Asp Pro Leu Pro Ile Ile Tyr Tyr Val Gly Arg Asn Val
275 280 285
Arg Val Glu Gln Leu Ser Asn Met Val Val Arg Ala Cys Lys Cys Ser
290 295 300

<210> 27

<211> 383

<212> PRT

<213> Rana sp.

<400> 27

Met Glu Val Leu Trp Met Leu Leu Val Leu Leu Val Leu His Leu Ser
1 5 10 15
Ser Leu Ala Met Ser Leu Ser Thr Cys Lys Ala Val Asp Met Glu Glu
20 25 30
Val Arg Lys Arg Arg Ile Glu Ala Ile Arg Gly Gln Ile Leu Ser Lys
35 40 45

Leu Lys Leu Asp Lys Ile Pro Asp Val Asp Ser Glu Lys Met Thr Val
 50 55 60
 Pro Ser Glu Ala Ile Phe Leu Tyr Asn Ser Thr Leu Glu Val Ile Arg
 65 70 75 80
 Glu Lys Ala Thr Arg Glu Glu Glu Glu His Val Gly His Asp Gln Asn
 85 90 95
 Ile Gln Asp Tyr Tyr Ala Lys Gln Val Tyr Arg Phe Glu Ser Ile Thr
 100 105 110
 Glu Leu Glu Asp His Glu Phe Lys Phe Lys Phe Asn Ala Ser Asn Val
 115 120 125
 Arg Glu Asn Val Gly Met Asn Ser Leu Leu His His Ala Glu Leu Arg
 130 135 140
 Met Tyr Lys Lys Gln Thr Asp Lys Asn Met Asp Gln Arg Met Glu Leu
 145 150 155 160
 Phe Trp Lys Tyr Gln Glu Asn Gly Thr Thr His Ser Arg Tyr Leu Glu
 165 170 175
 Ser Lys Tyr Ile Thr Pro Val Thr Asp Glu Trp Met Ser Phe Asp
 180 185 190
 Val Thr Lys Thr Val Asn Glu Trp Leu Lys Arg Ala Glu Glu Asn Glu
 195 200 205
 Gln Phe Gly Leu Gln Pro Ala Cys Lys Cys Pro Thr Pro Gln Ala Lys
 210 215 220
 Asp Ile Asp Ile Glu Gly Phe Pro Ala Leu Arg Gly Asp Leu Ala Ser
 225 230 235 240
 Leu Ser Ser Lys Glu Asn Thr Lys Pro Tyr Leu Met Ile Thr Ser His
 245 250 255
 Pro Ala Glu Arg Ile Asp Thr Val Thr Ser Ser Arg Lys Lys Arg Gly
 260 265 270
 Val Gly Gln Glu Tyr Cys Phe Gly Asn Asn Gly Pro Asn Cys Cys Val
 275 280 285
 Lys Pro Leu Tyr Ile Asn Phe Arg Lys Asp Leu Gly Trp Lys Trp Ile
 290 295 300
 His Glu Pro Lys Gly Tyr Glu Ala Asn Tyr Cys Leu Gly Asn Cys Pro
 305 310 315 320
 Tyr Ile Trp Ser Met Asp Thr Gln Tyr Ser Lys Val Leu Ser Leu Tyr
 325 330 335
 Asn Gln Asn Asn Pro Gly Ala Ser Ile Ser Pro Cys Cys Val Pro Asp
 340 345 350
 Val Leu Glu Pro Leu Pro Ile Ile Tyr Tyr Val Gly Arg Ile Ala Lys
 355 360 365
 Val Glu Gln Leu Ser Asn Met Val Val Arg Ser Cys Asn Cys Ser
 370 375 380

<210> 28

<211> 8

<212> PRT

<213> Homo sapiens

<400> 28

Ala Pro Gln Gly Ile Ala Gly Gln
1 5

<210> 29

<211> 8

<212> PRT

<213> Homo sapiens

<400> 29

Gly Pro Gln Gly Leu Leu Gly Ala
1 5

<210> 30

<211> 8

<212> PRT

<213> Homo sapiens

<400> 30

Gly Pro Gln Gly Leu Ala Gly Gln
1 5

<210> 31

<211> 8

<212> PRT

<213> Homo sapiens

<400> 31

Gly Pro Leu Gly Ile Ala Gly Ile
1 5

<210> 32

<211> 8

<212> PRT

<213> Homo sapiens

<400> 32

Gly Pro Glu Gly Leu Arg Val Gly
1 5

<210> 33

<211> 8

<212> PRT

<213> Rattus sp.

<400> 33

Ala Ala Tyr His Leu Val Ser Gln
1 5

<210> 34

<211> 8

<212> PRT

<213> Rattus sp.

<400> 34

Met Asp Ala Phe Leu Glu Ser Ser
1 5

<210> 35

<211> 8

<212> PRT

<213> Rattus sp.

<400> 35

Glu Pro Gln Ala Leu Ala Met Ser
1 5

<210> 36

<211> 8

<212> PRT

<213> Rattus sp.

<400> 36

Gln Ala Leu Ala Met Ser Ala Ile
1 5

<210> 37

<211> 8

<212> PRT

<213> Gallus domesticus

<400> 37

Pro Ser Tyr Phe Leu Asn Ala Gly
1 5

<210> 38

<211> 8

<212> PRT

<213> Homo sapiens

<400> 38

Tyr Glu Ala Gly Leu Gly Val Val
1 5

<210> 39

<211> 8

<212> PRT

<213> Homo sapiens

<400> 39

Ala Gly Leu Gly Val Val Glu Arg
1 5

<210> 40

<211> 8

<212> PRT

<213> Homo sapiens

<400> 40

Ala Gly Leu Gly Ile Ser Ser Thr
1 5

<210> 41

<211> 8

<212> PRT

<213> Unknown

<220>

<223> Sequence source uncertain

<400> 41

Gly Ala Met Phe Leu Glu Ala Ile
1 5

<210> 42

<211> 8

<212> PRT

<213> Homo sapiens

<400> 42

Ile Pro Glu Asn Phe Phe Gly Val
1 5

<210> 43

<211> 8

<212> PRT

<213> Homo sapiens

<400> 43

Thr Glu Gly Glu Ala Arg Gly Ser
1 5

<210> 44

<211> 8

<212> PRT

<213> Homo sapiens

<400> 44

Arg Ala Ile His Ile Gln Ala Glu
1 5

<210> 45

<211> 8

<212> PRT

<213> Homo sapiens

<400> 45

Leu Arg Ala Tyr Leu Leu Pro Ala
1 5

<210> 46

<211> 8

<212> PRT

<213> Cavia porcellus

<220>

<221> SITE

<222> (3)..(3)

<223> Xaa=Hyp

<400> 46

Gly Ala Xaa Gly Leu Glx Gly His
1 5

<210> 47

<211> 8

<212> PRT

<213> Rattus sp.

<400> 47

Gly Pro Gln Gly Val Arg Gly Glu
1 5

<210> 48

<211> 8

<212> PRT

<213> Rattus sp.

<400> 48

Gly Pro Ala Gly Val Gln Gly Pro
1 5

<210> 49

<211> 8
<212> PRT
<213> Rattus sp.

<220>
<221> SITE
<222> (6)..(6)
<223> Xaa=Hyp

<400> 49
Gly Pro Ser Gly Leu Xaa Gly Pro
1 5
<210> 50
<211> 8
<212> PRT
<213> Rattus sp.

<400> 50
Gly Pro Ala Gly Glu Arg Gly Ser
1 5
<210> 51
<211> 8
<212> PRT
<213> Rattus sp.

<400> 51
Gly Ala Lys Gly Leu Thr Gly Ser
1 5
<210> 52
<211> 8
<212> PRT
<213> Rattus sp.

<400> 52
Gly Pro Ala Gly Gln Asp Gly Pro
1 5
<210> 53
<211> 8
<212> PRT
<213> Rattus sp.

<400> 53

Gly Pro Ala Gly Phe Ala Gly Pro
1 5

<210> 54

<211> 8

<212> PRT

<213> Rattus sp.

<400> 54

Gly Pro Ile Gly Asn Val Gly Ala
1 5

<210> 55

<211> 8

<212> PRT

<213> Rattus sp.

<220>

<221> SITE

<222> (3)..(3)

<223> Xaa=Hyl

<400> 55

Gly Pro Xaa Gly Ser Arg Gly Ala
1 5

<210> 56

<211> 8

<212> PRT

<213> Bos taurus

<400> 56

Gly Pro Gln Gly Ile Ala Gly Gln
1 5

<210> 57

<211> 8

<212> PRT

<213> Bos taurus

<400> 57

Gly Pro Gln Gly Leu Leu Gly Ala
1 5

<210> 58

<211> 8

<212> PRT

<213> Homo sapiens

<400> 58

Ile Pro Glu Asn Phe Phe Gly Val
1 5

<210> 59

<211> 8

<212> PRT

<213> Homo sapiens

<400> 59

Pro Pro Gly Ala Tyr His Gly Ala
1 5

<210> 60

<211> 8

<212> PRT

<213> Homo sapiens

<400> 60

Arg Ala Ile His Ile Gln Ala Glu
1 5

<210> 61

<211> 8

<212> PRT

<213> Homo sapiens

<400> 61

Gly Pro His Leu Leu Val Glu Ala
1 5

<210> 62

<211> 8

<212> PRT

<213> Homo sapiens

<400> 62

Leu Arg Ala Tyr Leu Leu Pro Ala
1 5

<210> 63

<211> 8

<212> PRT

<213> Homo sapiens

<400> 63

Gly Pro Glu Gly Leu Arg Val Gly
1 5

<210> 64

<211> 8

<212> PRT

<213> Homo sapiens

<400> 64

Arg Val Gly Phe Tyr Glu Ser Asp
1 5

<210> 65

<211> 8

<212> PRT

<213> Homo sapiens

<400> 65

Leu Leu Ser Ala Leu Val Glu Thr
1 5

<210> 66

<211> 8

<212> PRT

<213> Unknown

<220>

<223> Sequence source uncertain

<400> 66

Glu Ala Ile Pro Met Ser Ile Pro
1 5

<210> 67

<211> 8

<212> PRT

<213> Unknown

<220>

<223> Sequence source uncertain

<400> 67

Ile Ala Gly Arg Ser Leu Asn Pro
1 5

<210> 68

<211> 8

<212> PRT

<213> Gallus domesticus

<400> 68

Leu Asn Ala Gly Phe Thr Ala Ser
1 5

<210> 69

<211> 8

<212> PRT

<213> Homo sapiens

<400> 69

Ile Pro Glu Asn Phe Phe Gly Val
1 5

<210> 70

<211> 8

<212> PRT

<213> Unknown

<220>

<223> Sequence source uncertain

<400> 70

Lys Pro Gln Gln Phe Phe Gly Leu
1 5

<210> 71

<211> 8

<212> PRT

<213> Homo sapiens

<400> 71

Asp Val Ala Gln Phe Val Leu Thr
1 5

<210> 72

<211> 8

<212> PRT

<213> Homo sapiens

<400> 72

Asp Thr Leu Glu Val Met Arg Lys
1 5

<210> 73

<211> 8

<212> PRT

<213> Homo sapiens

<400> 73

Asp Val Gly His Phe Arg Thr Phe
1 5

<210> 74

<211> 8

<212> PRT

<213> Homo sapiens

<400> 74

Asp Ser Gly Gly Phe Met Leu Thr
1 5

<210> 75

<211> 8

<212> PRT

<213> Homo sapiens

<400> 75

Arg Val Ala Glu Met Arg Gly Glu
1 5

<210> 76

<211> 8

<212> PRT

<213> Homo sapiens

<400> 76

Asp Leu Gly Arg Phe Gln Thr Phe
1 5

<210> 77

<211> 8

<212> PRT

<213> Homo sapiens

<400> 77

Pro Phe Ser Pro Leu Val Ala Thr
1 5

<210> 78

<211> 8

<212> PRT

<213> Homo sapiens

<400> 78

Leu Arg Ala Tyr Leu Leu Pro Ala
1 5

<210> 79

<211> 8

<212> PRT

<213> Unknown

<220>

<223> Sequence source uncertain

<400> 79

Ala Pro Gly Asn Ala Ser Glu Ser
1 5

<210> 80

<211> 8

<212> PRT

<213> Unknown

<220>

<223> Sequence source uncertain

<400> 80

Phe Ser Ser Glu Ser Lys Arg Glu
1 5

<210> 81

<211> 8

<212> PRT

<213> Bos taurus

<400> 81

Ala Gly Gly Ala Gln Met Gly Val
1 5

<210> 82

<211> 8

<212> PRT

<213> Bos taurus

<400> 82

Gln Met Gly Val Met Gln Gly Pro
1 5

<210> 83

<211> 8

<212> PRT

<213> Bos taurus

<400> 83

Met Ala Ala Ser Leu Lys Arg Pro
1 5

<210> 84

<211> 8

<212> PRT

<213> Bos taurus

<400> 84

Met Ala Ala Ser Ala Lys Arg Glu
1 5

<210> 85

<211> 8

<212> PRT

<213> Bos taurus

<400> 85

Met Ala Ala Ser Leu Arg Lys Pro
1 5

<210> 86

<211> 8

<212> PRT

<213> Bos taurus

<400> 86

Gln Ala Gln Ala Ile Leu Gln Gln
1 5

<210> 87

<211> 8

<212> PRT

<213> Homo sapiens

<400> 87

Arg Ala Ile His Ile Gln Ala Glu
1 5

<210> 88

<211> 8

<212> PRT

<213> Bos taurus

<400> 88

Leu Val Glu Ala Leu Tyr Leu Val
1 5

<210> 89

<211> 8

<212> PRT

<213> Bos taurus

<400> 89

Glu Ala Leu Tyr Leu Val Cys Gly
1 5

<210> 90

<211> 8

<212> PRT

<213> Homo sapiens

<400> 90

Ile Pro Glu Asn Phe Phe Gly Val
1 5

<210> 91

<211> 8

<212> PRT

<213> Homo sapiens

<400> 91

Gly Pro His Leu Leu Val Glu Ala
1 5

<210> 92

<211> 8

<212> PRT

<213> Homo sapiens

<400> 92

Pro Pro Glu Glu Leu Lys Phe Gln
1 5

<210> 93

<211> 8

<212> PRT

<213> Homo sapiens

<400> 93

Gly Pro Pro Gly Val Val Gly Pro
1 5

<210> 94

<211> 8

<212> PRT

<213> Homo sapiens

<400> 94

Gly Pro Pro Gly Leu Arg Gly Glu
1 5

<210> 95

<211> 8

<212> PRT

<213> Homo sapiens

<400> 95

Gly Pro Glu Gly Val Val Gly Pro
1 5

<210> 96

<211> 8

<212> PRT

<213> Homo sapiens

<400> 96

Ile Pro Glu Asn Phe Phe Gly Val
1 5

<210> 97

<211> 8

<212> PRT

<213> Homo sapiens

<400> 97

Pro Pro Gly Ala Tyr His Gly Ala
1 5

<210> 98

<211> 8

<212> PRT

<213> Homo sapiens

<400> 98

Arg Ala Ile His Ile Gln Ala Glu
1 5

<210> 99

<211> 8

<212> PRT

<213> Homo sapiens

<400> 99

Arg Ala Ile His Ile Gln Ala Glu
1 5

<210> 100

<211> 8

<212> PRT

<213> Homo sapiens

<400> 100

Gly Pro His Leu Leu Val Glu Ala
1 5